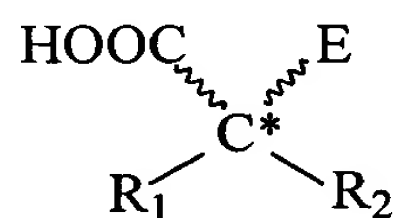

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19. The enantiomerically pure α -substituted carboxylic acid according to claim 18, having the structure:



R₁ and R₂ are each independently -H, substituted or unsubstituted alkyl, alkenyl, alkynyl, aryl, heteroaryl, cycloalkyl, heterocyclic, wherein said substituents are lower alkyl, hydroxy, alkoxy, mercapto, cycloalkyl, heterocyclic, aryl, heteroaryl, aryloxy, or halogen or optionally R₁ and R₂ are linked to cooperate to form a functional cyclic moiety, and

E is $-N(R_x)_2$ or $-OH$, wherein each R_x is $-H$ or lower alkyl.

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21. The enantiomerically pure α -substituted carboxylic acid according to claim 18, wherein the carboxylic acid is an α -hydroxy acid.

22. The method according to claim 1, wherein the nitrilase has an amino acid sequence as set forth in SEQ ID NO:2 or SEQ ID NO:4.

23. The method according to claim 1, wherein the nitrilase is encoded by a nucleic acid sequence as set forth in SEQ ID NO:1 or SEQ ID NO:3.

24. The method according to claim 1, wherein the nitrilase has an amino acid sequence at least 70% identical to the amino acid sequence as set forth in SEQ ID NO:2 or SEQ ID NO:4 and has nitrilase activity.

5 / 25. A substantially purified polypeptide having an amino acid sequence as set forth in SEQ ID NO:2 or SEQ ID NO:4 and sequences having at least 70% identity thereto and having nitrilase activity.

10 / 26. An isolated nucleic acid sequence encoding an amino acid sequence as set forth in SEQ ID NO:2 or SEQ ID NO:4 and sequences having at least 70% identity thereto and having nitrilase activity, and fragments thereof that hybridize to the nucleic acid sequence.

15 / 27. An isolated nucleic acid sequence as set forth in SEQ ID NO:1.

/ 28. An isolated nucleic acid sequence as set forth in SEQ ID NO:3.

20 / 29. A substantially purified polypeptide having an amino acid sequence as set forth in SEQ ID NO:2.

/ 30. A substantially purified polypeptide having an amino acid sequence as set forth in SEQ ID NO:4.

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